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Title: Reconfiguring the metabolism of photosynthetic microbes for their development as biotechnological platforms

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Seminar- Cemvita Factory Inc.

Reconfiguring the metabolism of photosynthetic microbes for their development as biotechnological platforms



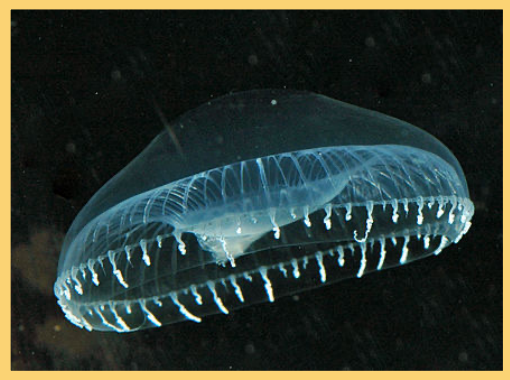
Raul Gonzalez

8/13/2021



Managed by Triad National Security, LLC for the U.S. Department of Energy's NNSA

Utilization of the marine biodiversity to foster technological innovation



Bioluminescence
(i.e. GFP)



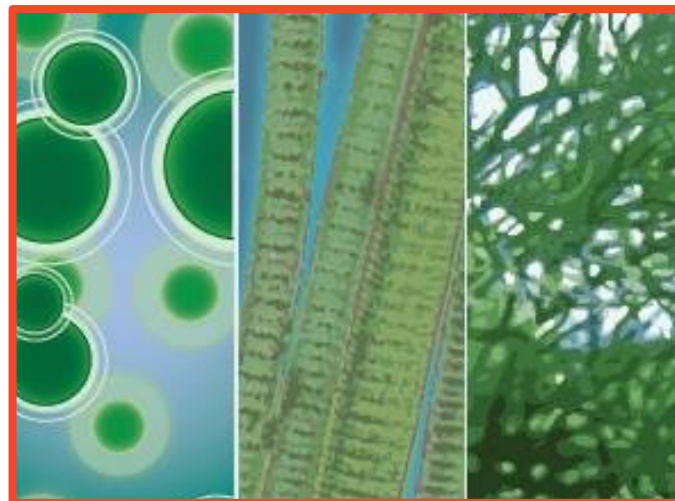
Improved fisheries
(i.e. Atlantic Salmon)



Bioremediation
(i.e. *Alcanivorax*)

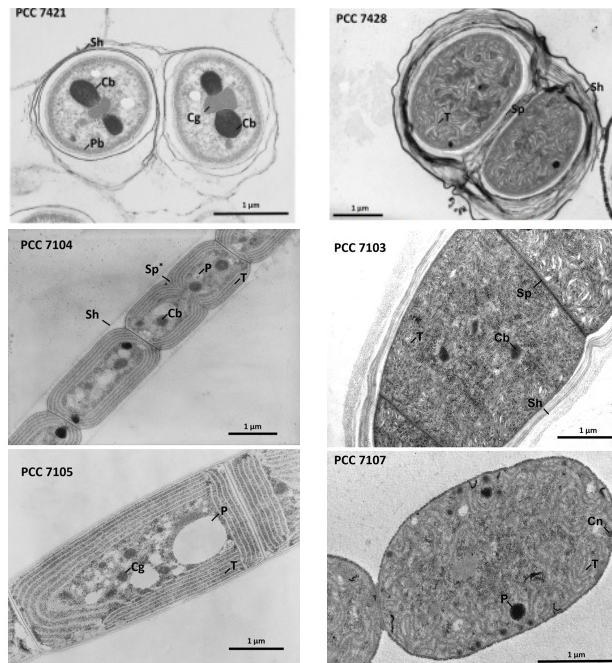


Bioactive compounds
(Limulus Amoebocyte
Lysate)

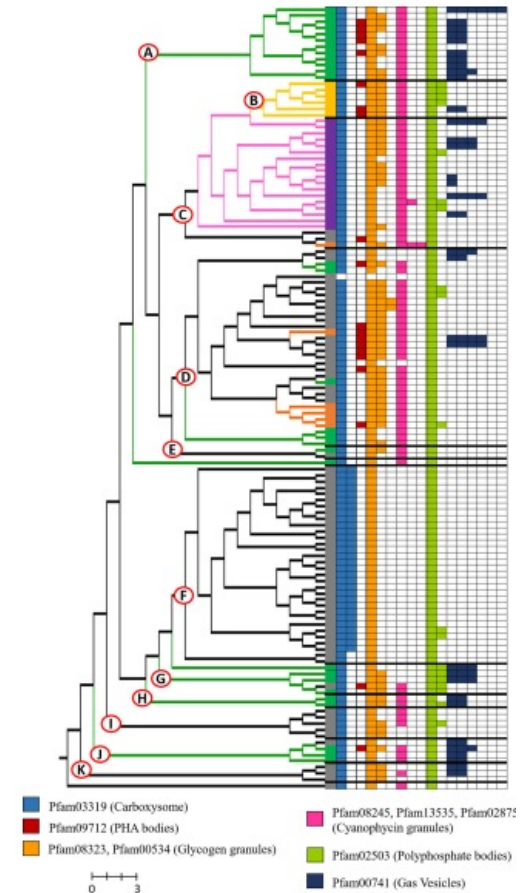
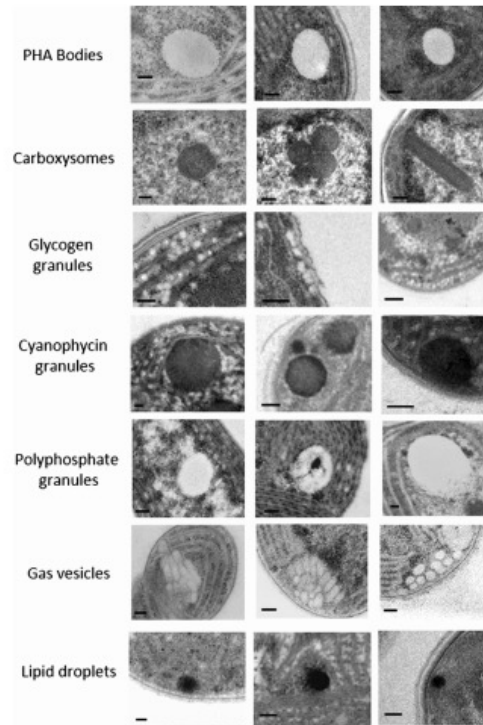


**Algal Biofuels
and Bioproducts**

Cyanobacterial ultrastructure and its relation to genomic features (Pfams)

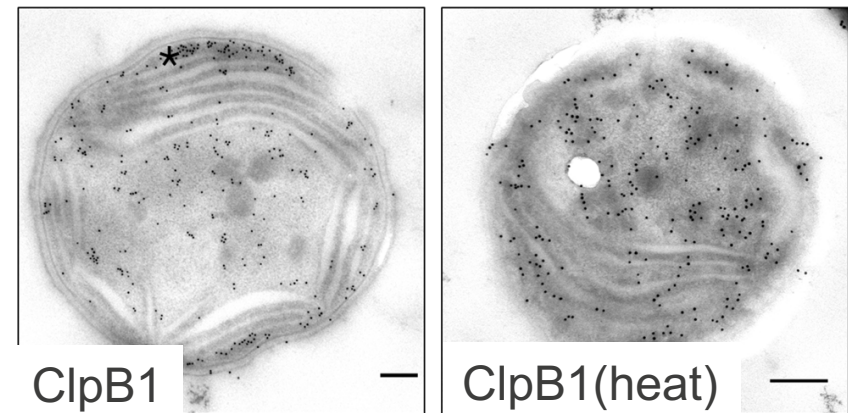
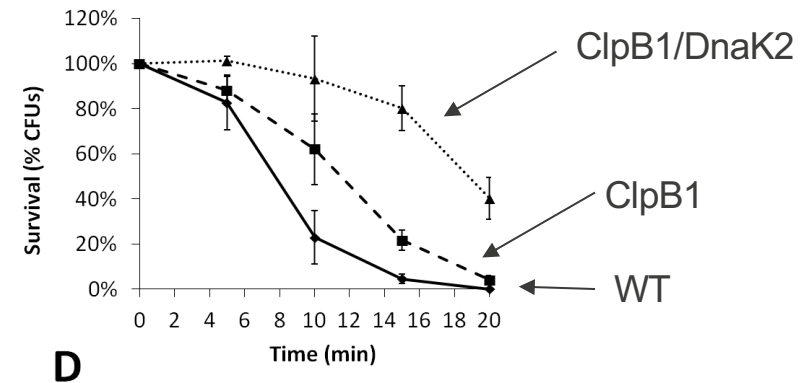
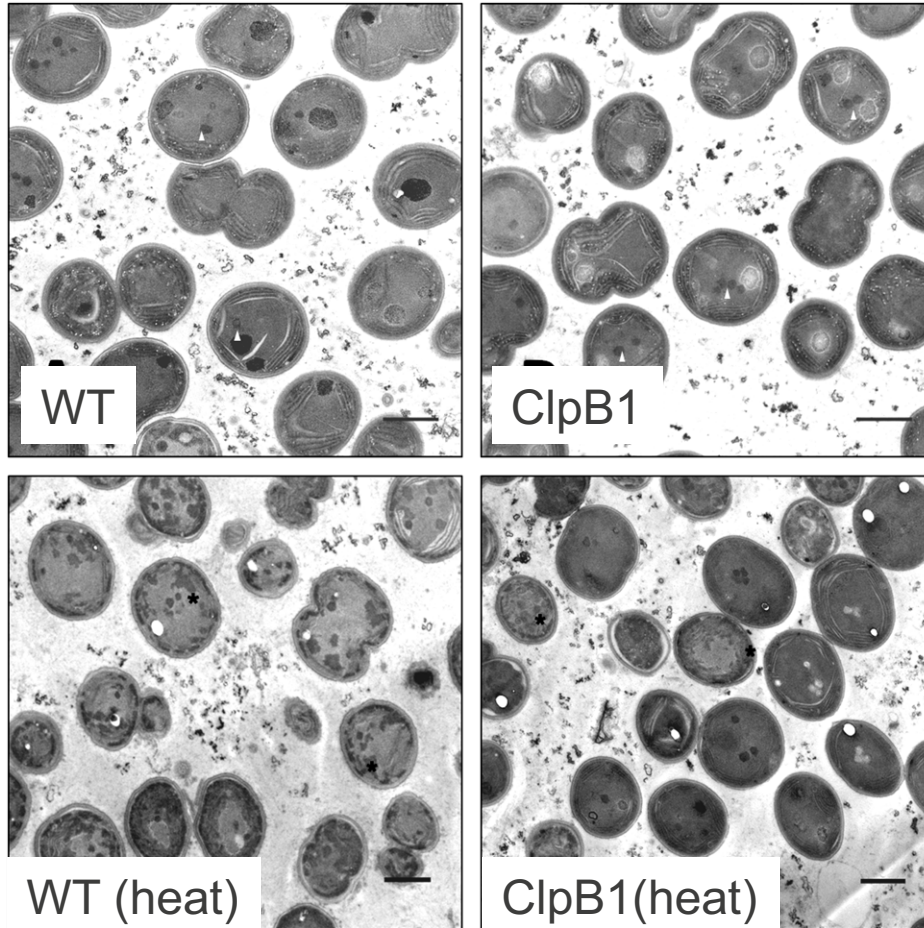


Photosynth Res (2016) 129:147–157



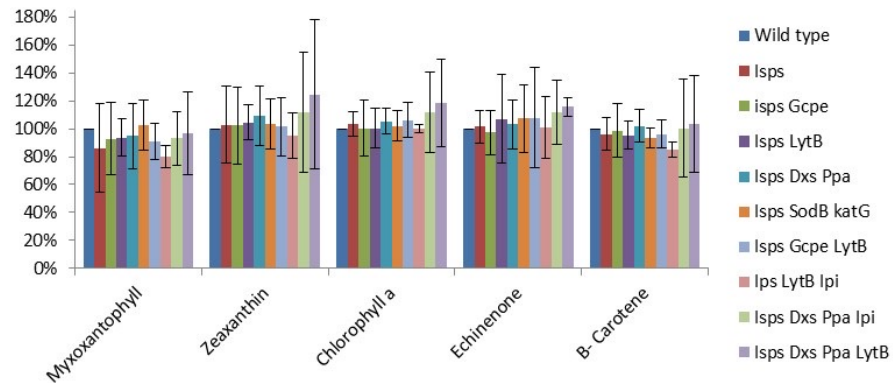
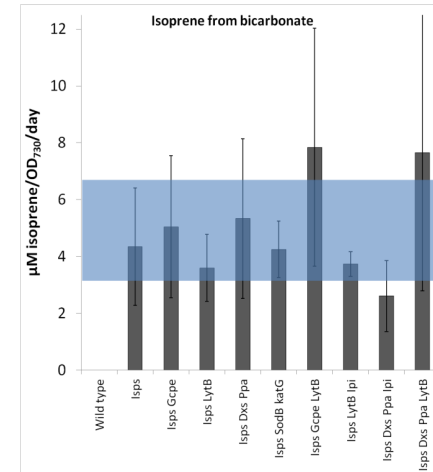
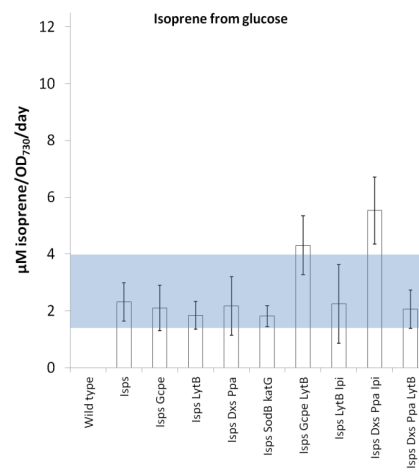
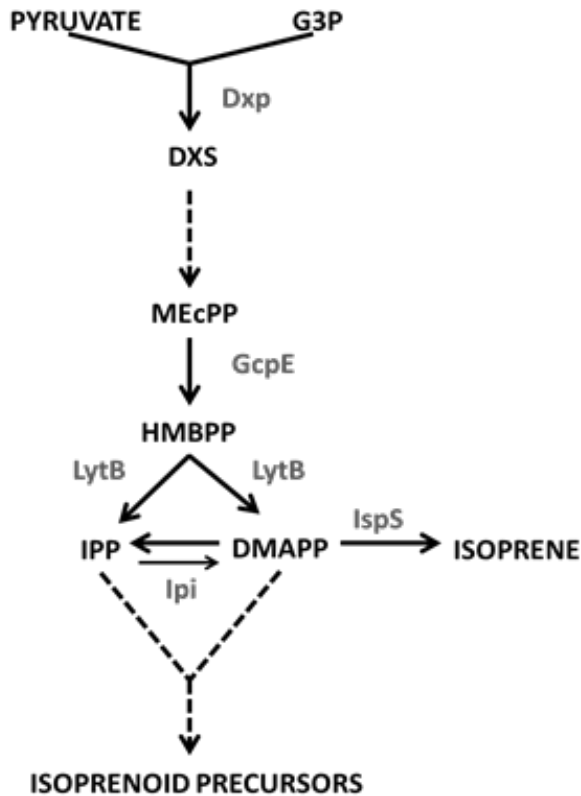
Improve environmental stress tolerance

Increase tolerance to heat stress as a proxy for other stresses



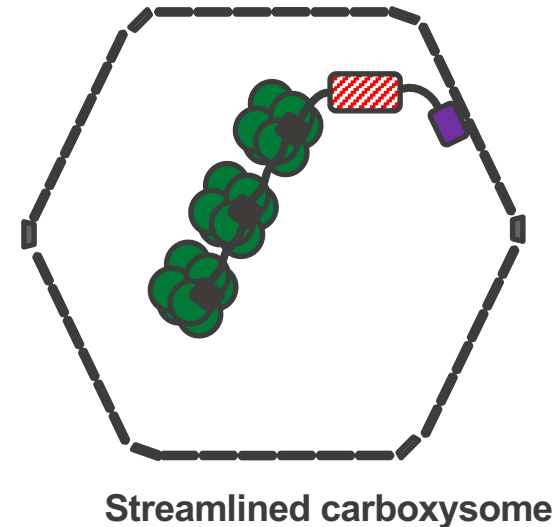
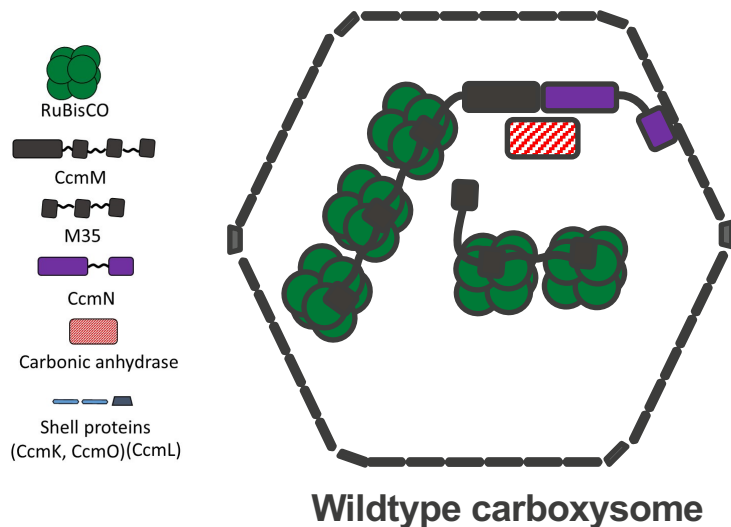
Increase Product Output

Increase yields of an isoprene-evolving mutant via pathway engineering



Improve CO₂ fixation machinery

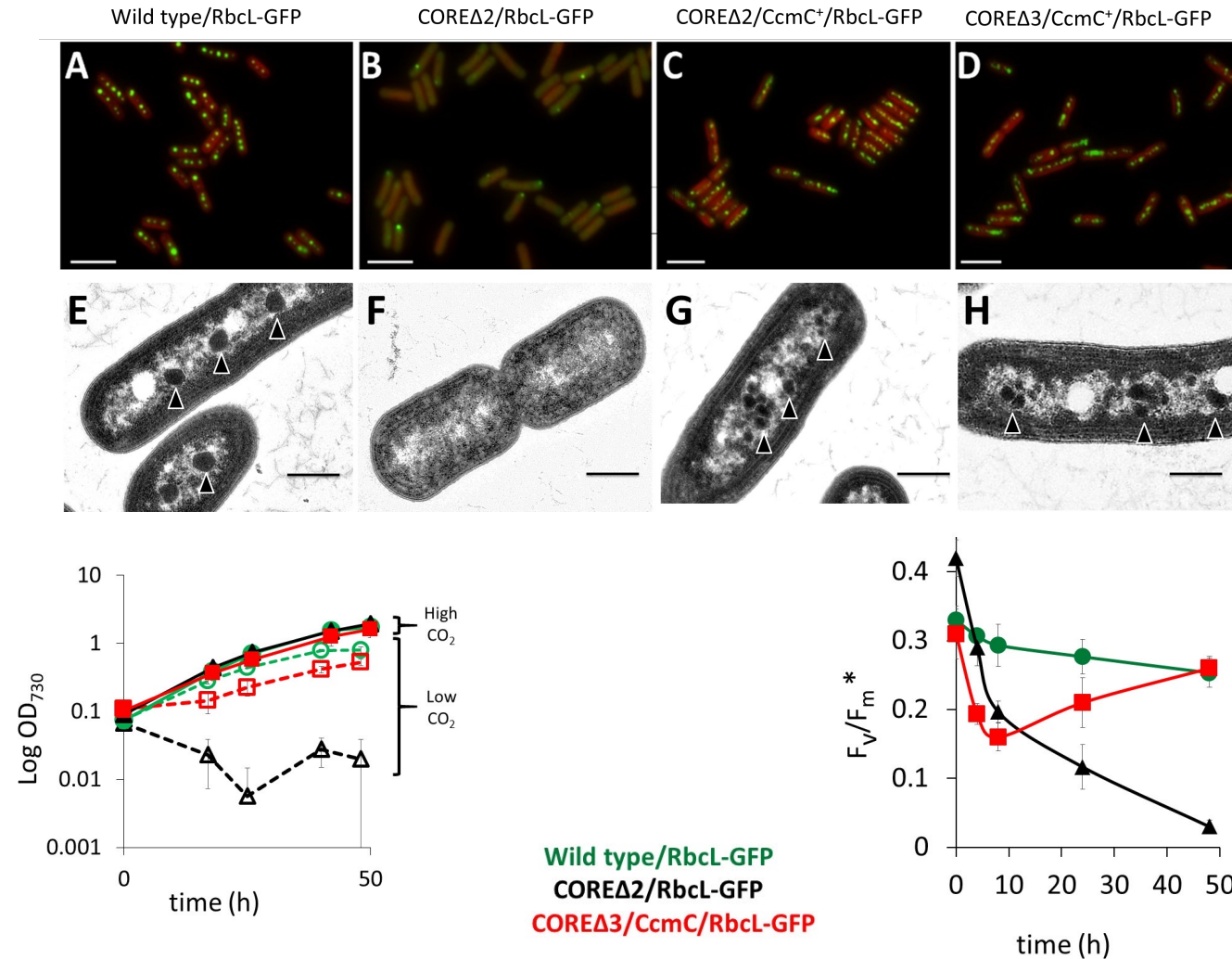
Streamlined assembly of carboxysomes



- Flexibility of subunits under environmental conditions
- Several subunits required for transfer to other organisms
- Complex assembly and regulation in other organisms
- Fixed stoichiometry of the composing domains
- Increased portability (fewer number of proteins required)
- Simplified assembly and regulation (one core assembly protein) in other organisms
- Potential for improvement

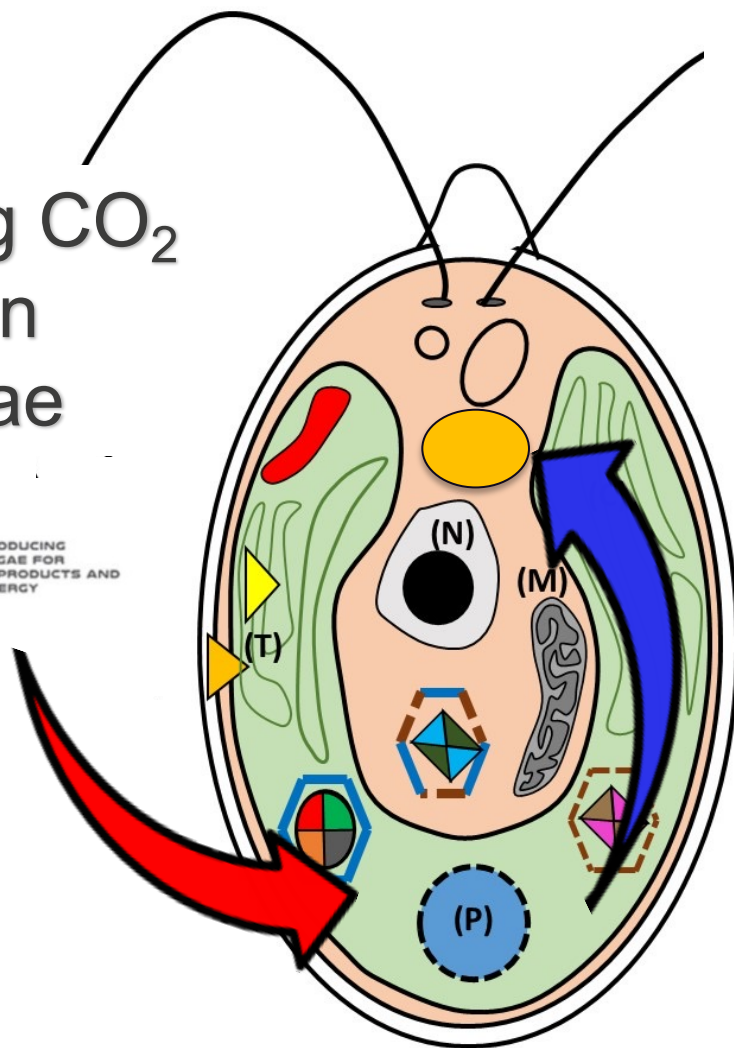
Improve CO₂ fixation machinery

Streamlined assembly of carboxysomes



Research at LANL

1) Increasing CO₂ fixation in microalgae



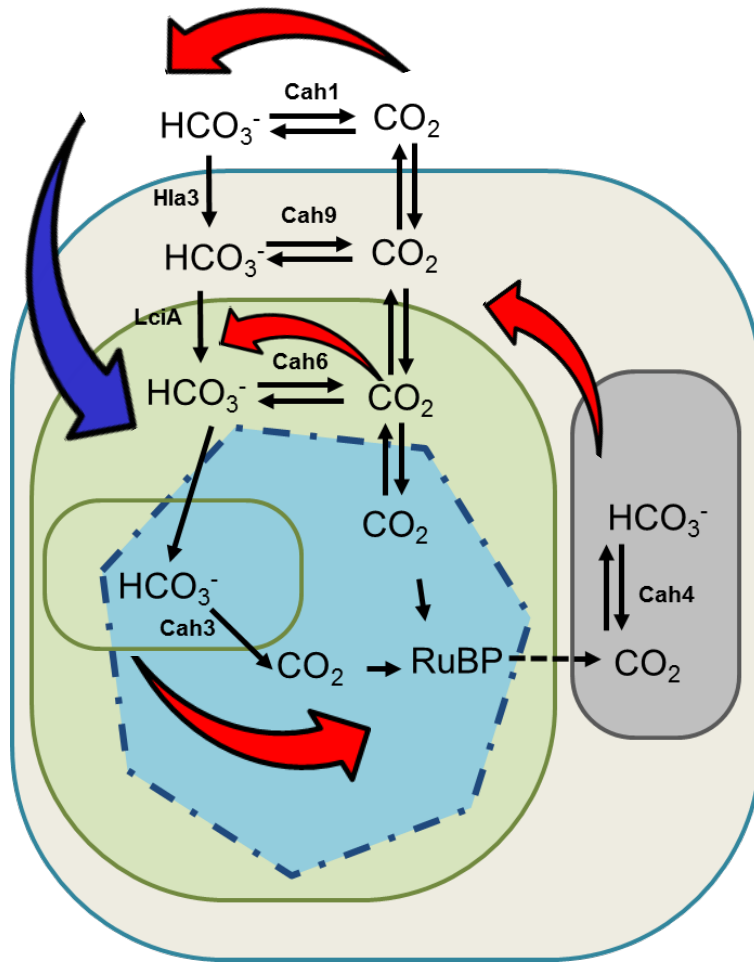
3) Customized PHA production in cyanobacteria



2) Increasing carbon storage compounds in microalgae

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Increasing carbon flux towards RuBisCO



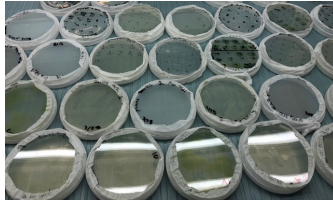
Carbonic anhydrase overexpression:

- **CAH1** and **CAH6**: Shift the equilibrium towards bicarbonate (decrease CO_2 leakage).
- **CAH3**: Produce more CO_2 available for RuBisCO.
- **CAH4**: Mobilize bicarbonate pool from mitochondria to the CCM.
- **BCA**: Shown previously to increase photosynthetic efficiency in plants.

HCO_3^- transporter overexpression:

- **HLA3** and **LCIA**: Increase the bicarbonate flux towards the chloroplast.

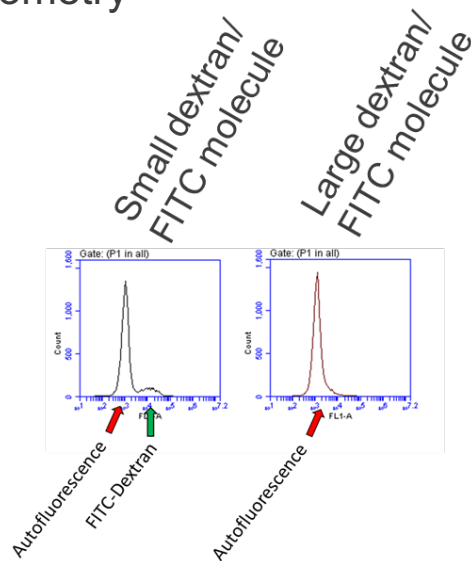
Challenge overcome: Development of a transformation method for *Chlorella sorokiniana*



- Biolistic DNA delivery and antibiotic selection frequently yielded too many false positives. Screening *practically* impossible.

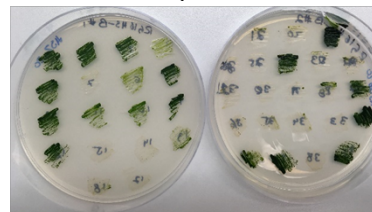
✓ Electroporation, Flow cytometry and Physiology

- Delivered fluorescent molecules by electroporation, screened by flow cytometry

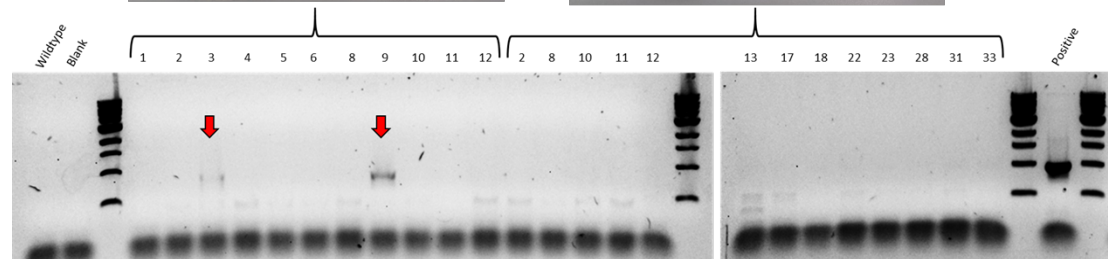
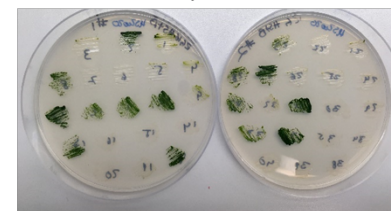


- Screened different culture conditions for recovery, selective pressure.

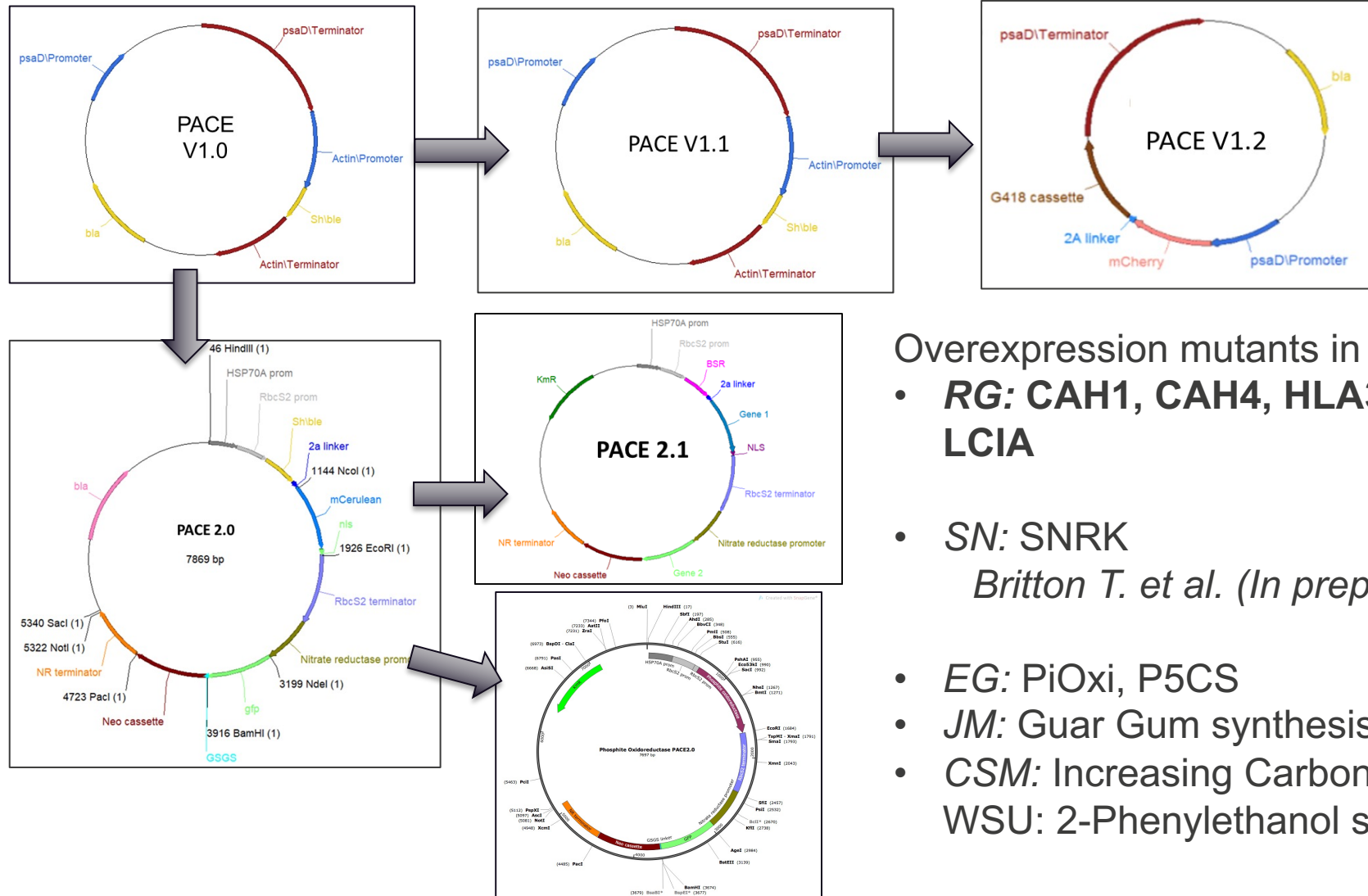
HS-B: 26/40 colonies



HS+B: 24/40 colonies



LANL team pushes boundaries of the state-of-technology



Overexpression mutants in CS:

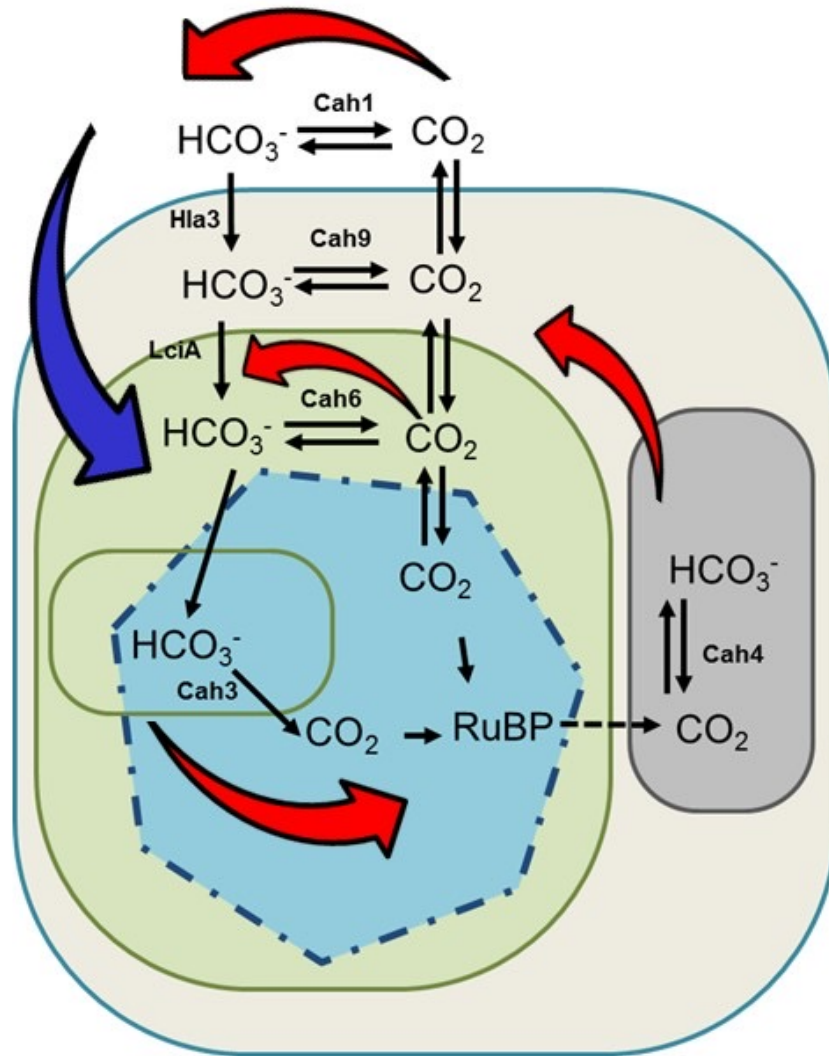
- **RG: CAH1, CAH4, HLA3, BCA, LCIA**

- **SN: SNRK**

Britton T. et al. (In preparation)

- **EG: PiOxi, P5CS**
- **JM: Guar Gum synthesis**
- **CSM: Increasing Carbon sink**
- **WSU: 2-Phenylethanol synthesis**

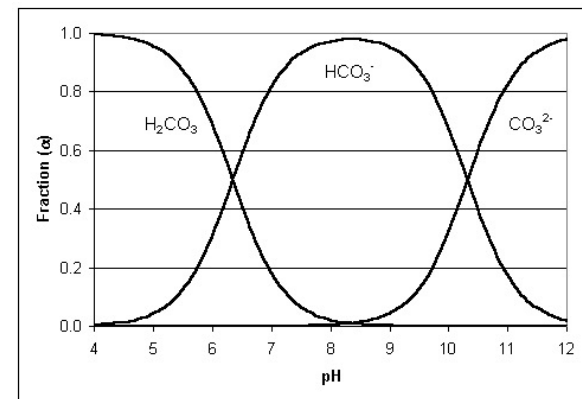
Gene stacking likely required to increase carbon fixation rates



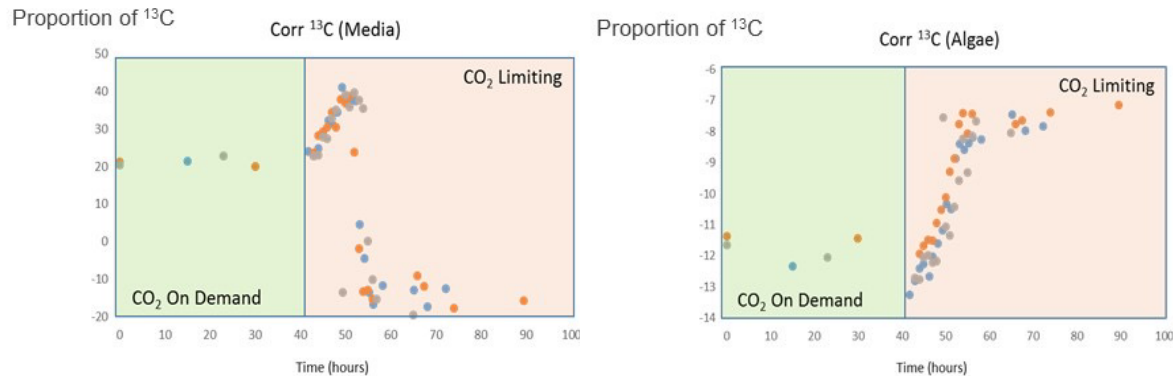
Single overexpression mutants generated in CS:

- **RG: CAH1, CAH4, HLA3, BCA, LCIA**

➤ No positive phenotype observed in single mutants. Likely, multiple gene overexpression required to cross all cellular barriers



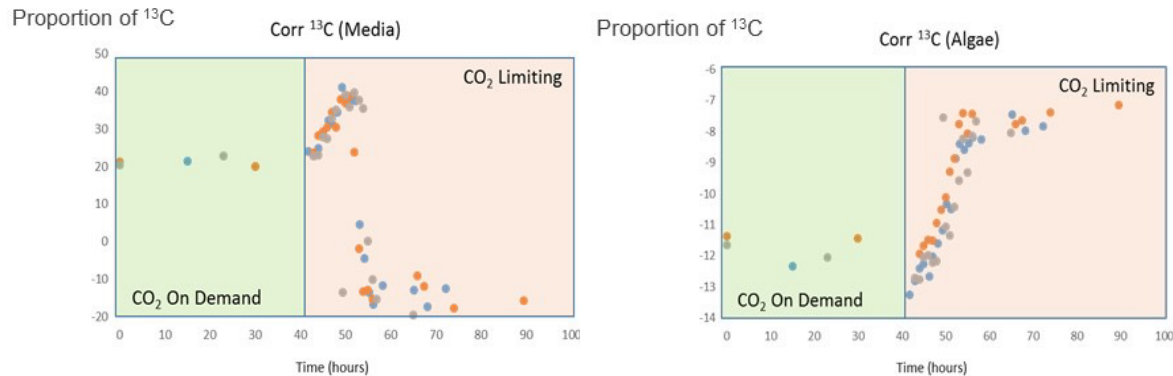
Photosynthetic capacity studied by limiting carbon availability



- ^{13}C discrimination tool developed in *N. salina* was implemented for measuring carbon use efficiency in *Chlorella sorokiniana*

Zidenga et al. *Algal Research* (2018)

Photosynthetic capacity studied by limiting carbon availability

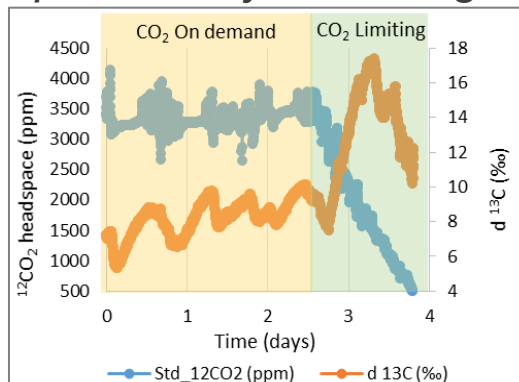


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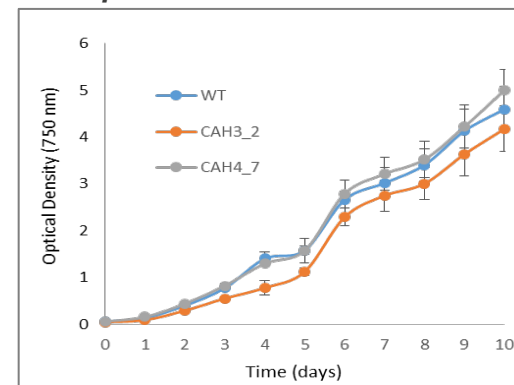
Zidenga et al. *Algal Research* (2018)

... which we are using to evaluate Carbonic anhydrase (CA) mutants in *Nannochloropsis gaditana* (Matt Posewitz, CSM)

Headspace analysis of N. gaditana

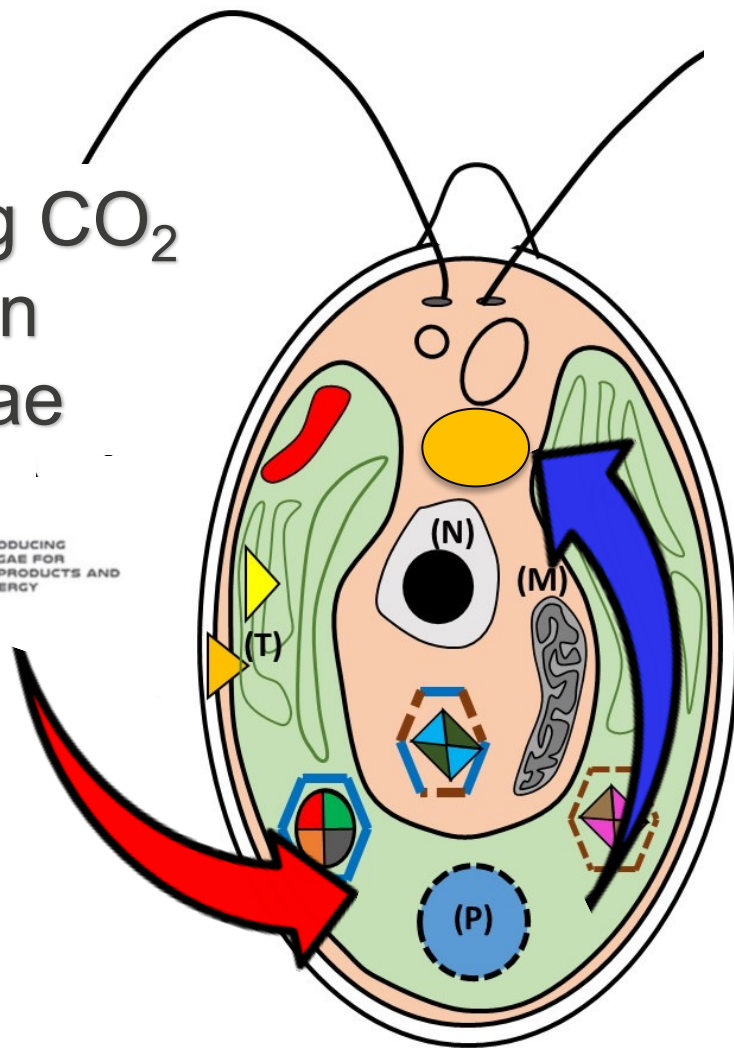


Growth comparison of CA mutants vs WT



Presentation Outline

1) Increasing CO₂ fixation in microalgae



3) Customized PHA production in cyanobacteria

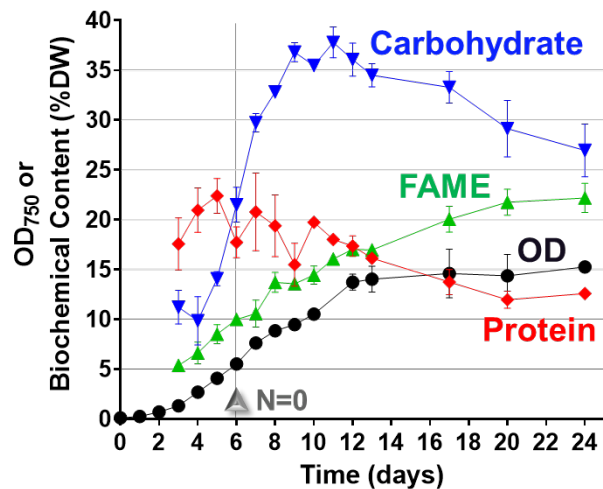


2) Increasing carbon storage compounds in microalgae

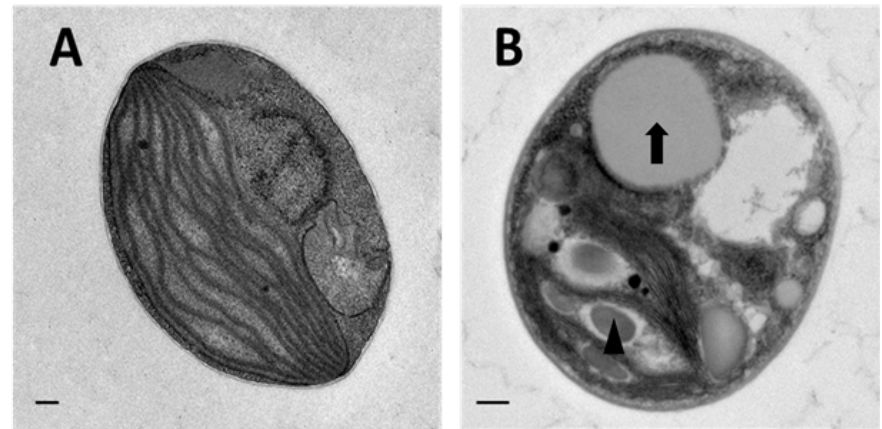
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Multiscale Characterization: Increasing carbon storage molecules in algal biomass

- Algae accumulate carbon storage compounds (lipids, carbohydrates) after Nitrogen depletion

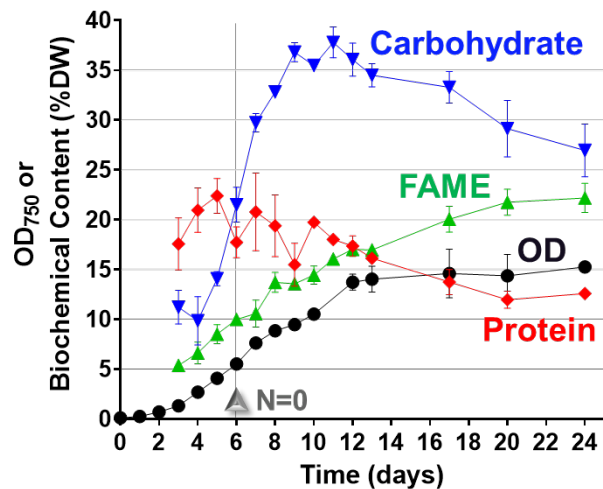


P. soloecismus

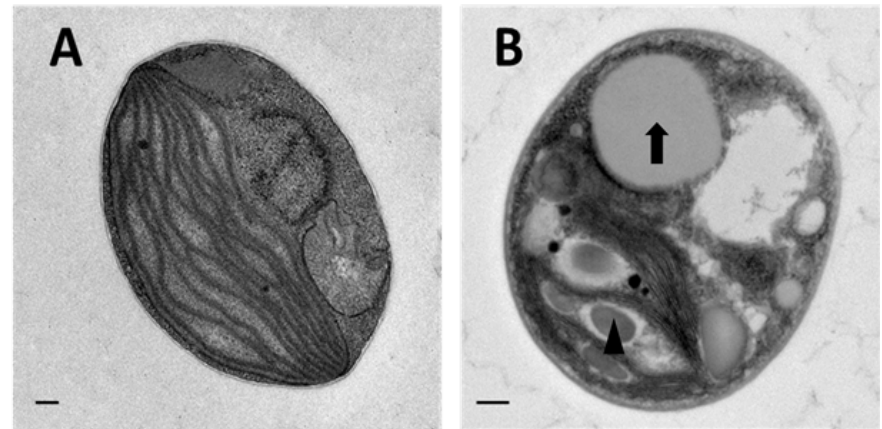


Multiscale Characterization: Increasing carbon storage molecules in algal biomass

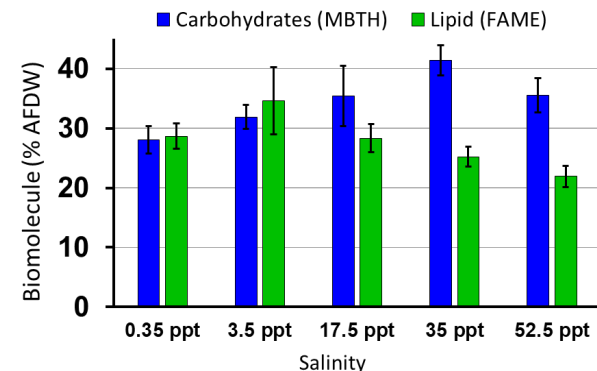
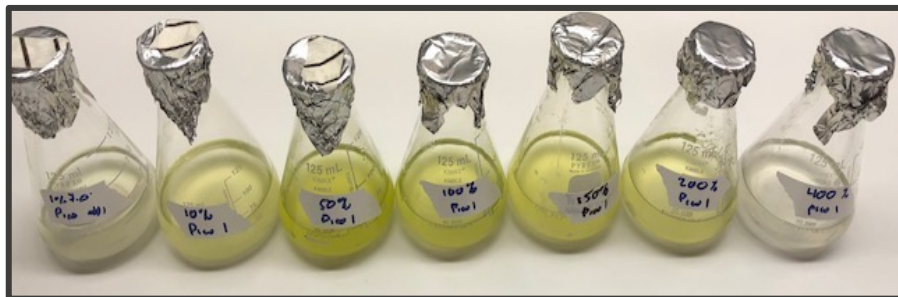
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P. soloecismus



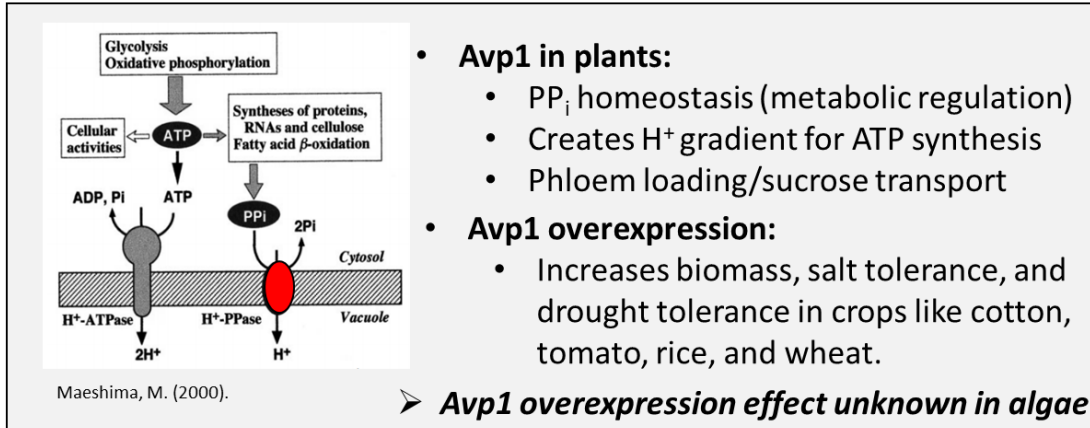
- Proportion of lipids vs carbohydrates may vary under culture conditions



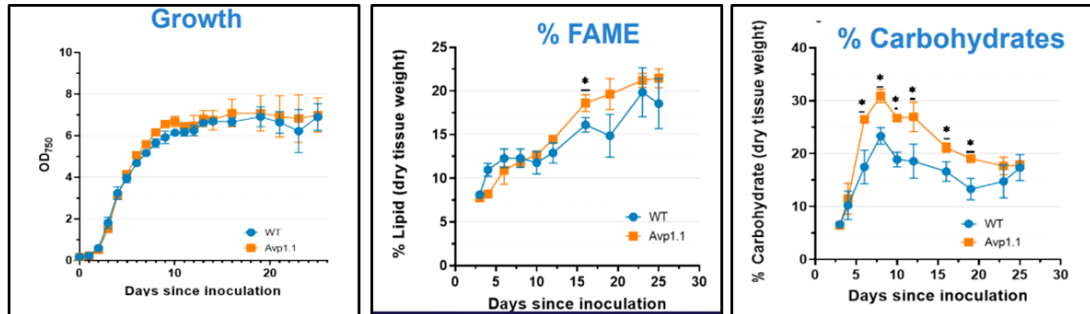
Gonzalez-Esquer CR et al. *Algal Research*

Increasing Carbon Storage in Transgenic Algae

Overexpression of a H⁺-pumping pyrophosphatase (AVP1)

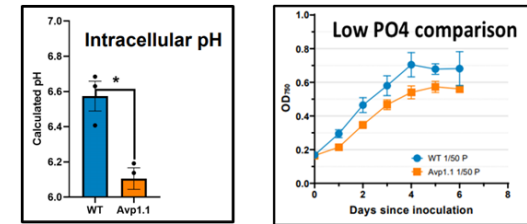


- AVP1 overexpression increases carbohydrates under N deplete conditions*

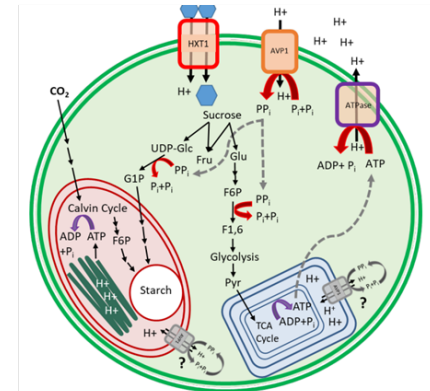


K Wright, R. Gonzalez, T. Dale

- AVP1 overexpression decreases intracellular pH and diminishes growth on low PO₄ conditions*



- AVP1 overexpression improves biochemical composition (increased C storage) of algal biomass, likely due to PP_i synthesis (provoking an intracellular P-limiting condition).*

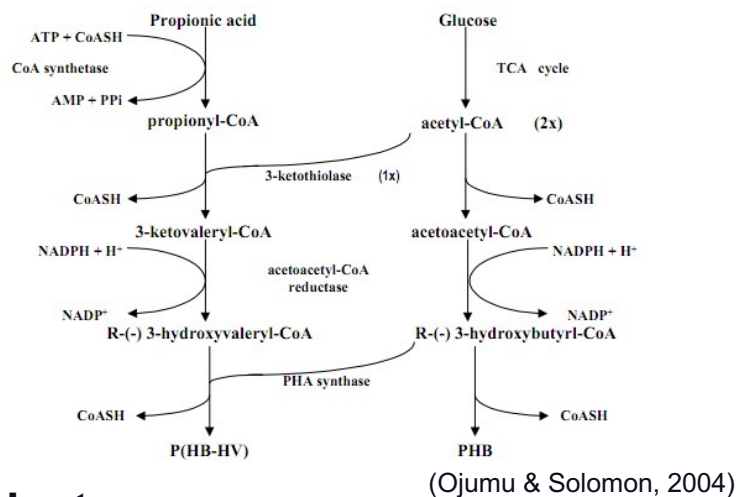


BioManiac: Can we leverage biodiversity to match/increase performance of chemically developed materials?

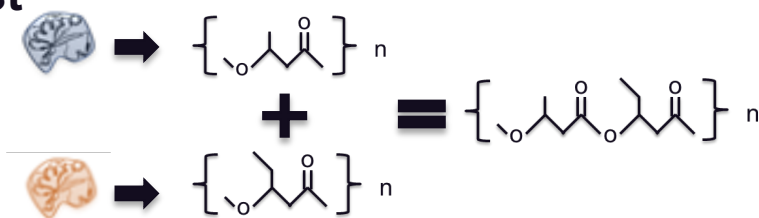
We use **Polyhydroxyalkanoates** (PHA) as case study. PHAs are carbon storage molecules with can be used as bioplastic precursors.

PHA cyanobacteria \neq PHA heterotrophic bacteria

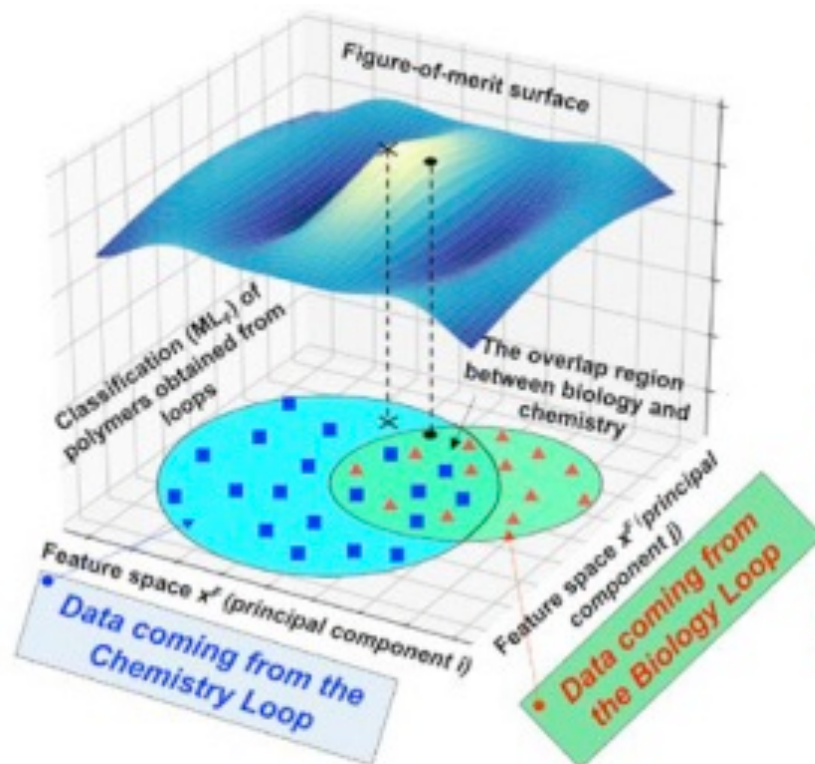
Metabolism



Biocatalyst

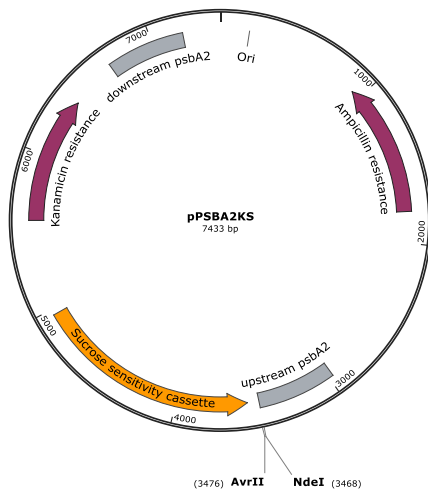


Machine Learning



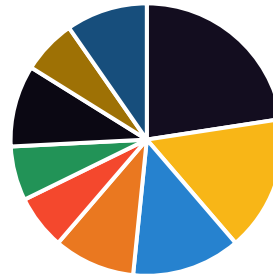
Assessment of ML-Predicted Genes by Overexpression

- Gene targets were synthesized into Psba2KS vector and transformed into PCC 6803

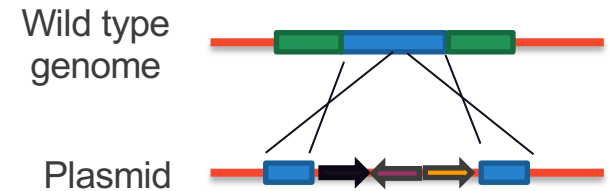


Synechocystis sp. PCC 6803 mutants (Total: 31)

- Unknown
- Signaling
- Lipid processes
- Acetate metabolism
- PHB synthesis
- Protein metabolism



- PCR screening of genomic integration



Testing mutant genome

Precise bp length; Wt negative

1)

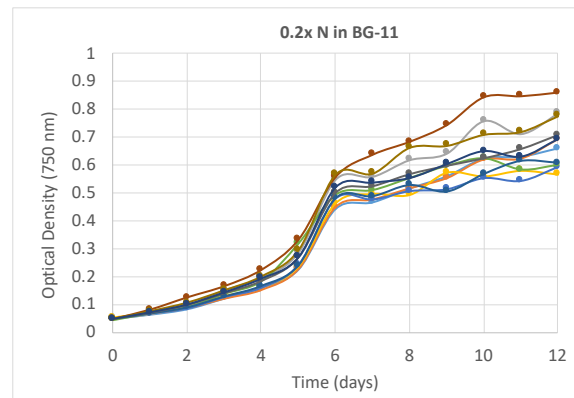
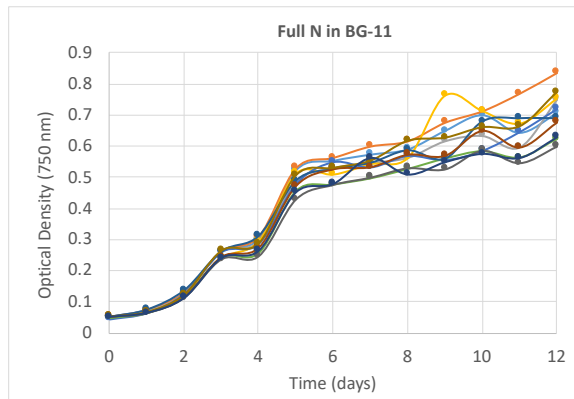


One band: segregated mutant; two bands: WT still present

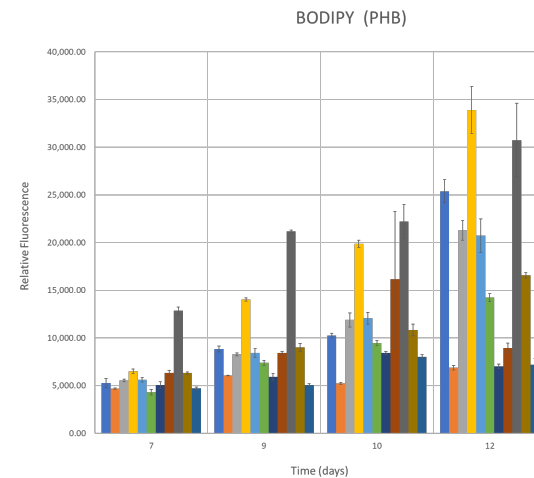
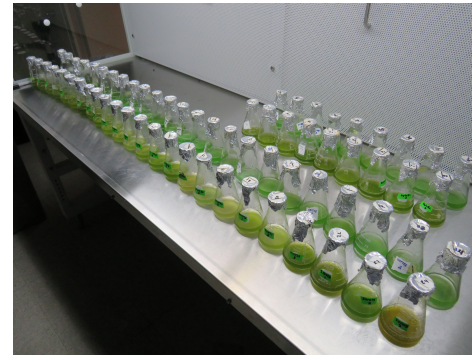


Validation (CyanoGEBA-ML in PCC 6803)

- Growth phenotypes in two different culture conditions



- PHB accumulation via BODIPY/Flow cytometry



Concluding remarks

- LANL is at the forefront of genetic engineering of microalgae
 - We have developed genetic engineering toolboxes for many strains, including *Picochlorum soloecismus*, *Nannochloropsis salina*, *Chlorella sorokiniana*
 - We have implemented such toolboxes for generating mutants with favorable phenotypes
- We have the opportunity to integrate metabolic features from different species (cyanobacteria>microalgae>plants) into a synthetic biology discovery and developmental platform
 - We can leverage the cyanobacterial metabolic “simplicity” to engineer complex organisms, i.e. for the production of renewable polymers
 - Unrivaled ‘omics and machine learning scientific collaboration